At p. 1, line 2, please insert as follows:

--This application is a continuation of U.S. Serial No. 09/052,521, filed March 30, 1998, which is a continuation-in-part of U.S. Serial No. 08/880,855, filed June 23, 1997, which is a continuation-in-part of U.S. Serial No. 08/842,842, filed April 16, 1997, now U.S. Patent No. 5,843,678, which are hereby incorporated by reference. --

At page 40, please amend the paragraph at lines 32-35 and page 41, lines 1-3, as follows:

This construct was engineered to be 242 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met(75)-Asp-Pro-Asn-Arg-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 41). The template to be used for PCR was pcDNA/32D-F3 and oligonucleotides #1581-72 and #1581-76 were the primer pair to be used for PCR and cloning this gene construct.

At page 41, please amend the paragraph at lines 13-19, as follows:

This construct was engineered to be 223 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-His(95)-Glu-Asn-Ala-Gly-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 44). The template used for PCR was pcDNA/32D-F3 and oligonucleotides #1591-90 and #1591-95 were the primer pair used for PCR and cloning this gene construct.

At page 41, please amend the paragraph at lines 29-35, as follows:

This construct was engineered to be 211 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-Ser(107)-Glu-Asp-Thr-Leu-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 45). The template used for PCR was pcDNA/32D-F3 and oligonucleotides #1591-93 and #1591-95 were the primer pair used for PCR and cloning this gene construct.

At page 42, please amend the paragraph at lines 11-17, as follows:

This construct was engineered to be 199 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met(118)-Lys-Gln-Ala-Phe-Gln-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 46). The template used for PCR was pcDNA/32D-F3 and oligonucleotides #1591-94 and #1591-95 were the primer pair used for PCR and cloning this gene construct.

C-terminal residues, NH_2-Met-Lys(128)-Glu-Leu-Gln-His-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 47). The

template used for PCR was pcDNA/32D-F3 and oligonucleotides #1591-91 and #1591-95 were the primer pair used for PCR and cloning this gene construct.

At page 43, please amend the paragraph at lines 5-11, as follows:

This construct was engineered to be 181 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-Gln(137)-Arg-Phe-Ser-Gly-------Gln-Asp-Ile-Asp(316)-COOH (SEQ ID NO: 48). The template used for PCR was pcDNA/32D-F3 and oligonucleotides #1591-92 and #1591-95 were the primer pair used for PCR and cloning this gene construct.

At page 43, please amend the paragraph at lines 21-28, as follows:

This construct is engineered to be 171 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met(146)-Glu-Gly-Ser-Trp--------Gln-Asp-Ile-Asp(316)- COOH (SEQ ID NO: 49). The template to be used for PCR is pAMG21-murine OPG binding protein [75-316] described above and oligonucleotides #1600-98 and #1581-76 will be the primer pair to be used for PCR and cloning this gene construct.

At page 44, please amend the paragraph at lines 2-9, as follows:

This construct is engineered to be 162 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-Arg(156)-Gly-Lys-Pro--------Gln-Asp-Ile-Asp(316)- COOH (SEQ ID NO: 50). The template to be used for PCR is pAMG21-murine OPG binding protein [158-316] below and oligonucleotides #1619-86 and #1581-76 will be the primer pair to be used for PCR and cloning this gene construct.

At page 44, please amend the paragraph at lines 19-25, as follows:

This construct was engineered to be 160 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-Lys(158)-Pro-Glu-Ala--------Gln-Asp-Ile-Asp(316)- COOH (SEQ ID NO: 51). The template to be used for PCR was pcDNA/32D-F3 and oligonucleotides #1581-73 and #1581-76 were the primer pair to be used for PCR and cloning this gene construct.

At page 44, please amend the paragraph at lines 35-37 and page 45, lines 1-4, as follows:

This construct is engineered to be 152 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-His(166)-Leu-Thr-Ile--------Gln-Asp-Ile-Asp(316)- COOH (SEQ ID NO: 52). The template to be used for PCR is pcDNA/32D-F3 and oligonucleotides #1581-75 and #1581-76 will be the primer pair

At page [illegible] paragraph at lines [illegible]

This construct is engineered to be 150 amino acids in length and have the following N-terminal and C-terminal residues, NH_2-Met-Thr(168)-Ile-Asn-Ala--------Gln-Asp-Ile-Asp(316)- COOH (SEQ ID NO: 53). The

template to be used for PCR is pcDNA/32D-F3 and oligonucleotides #1581-74 and #1581-76 will be the primer pair to be used for PCR and cloning.

At page 63, please amend the paragraph at line 13, as follows:

NH2 - K L V T L Q V T P-CO2H (SEQ ID NO: 54).